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                 alerts (SDIs) affected
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                COMPUAB reloaded; updating to resume; current-awareness
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NEWS 15 DEC 30 CAPLUS - PATENT COVERAGE EXPANDED
NEWS 16 JAN 03 No connect-hour charges in EPFULL during January and
                February 2005
NEWS EXPRESS OCTOBER 29 CURRENT WINDOWS VERSION IS V7.01A, CURRENT
             MACINTOSH VERSION IS V6.0c(ENG) AND V6.0Jc(JP),
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L2 11 WEVLCWTWETCER/SQSP

=> s wevlcwtwetcer/sqep

2 WEVLCWTWETCER/SQEP

309925 SQL=13

L3 2 WEVLCWTWETCER/SQEP (WEVLCWTWETCER/SQEP AND SQL=13)

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=> s {0,99}.wevlcwtwetcer.{0,99}./sqsp INVALID USE OF BRACE OPERATOR

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L8 6 WEVLCWTWETCER. {0,99}./SQSP

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L9 9 L6 OR L7 OR L8

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L11 ANSWER 1 OF 4 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 1

AN 2004:372929 CAPLUS

DN 140:395489

TI Sequences of blood-coagulation factor VIIa-binding peptides

IN Lazarus, Robert A.; Maun, Henry R.

PA Genentech, Inc., USA

SO U.S. Pat. Appl. Publ., 102 pp.

CODEN: USXXCO

DT Patent

LA English

FAN. CNT 1

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
PI US 2004087767	A1	20040506	US 2003-356257	20030130
PRAI US 2002-355420P	P	20020206		

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blood-coagulation factor VIIa-binding peptides)
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     RL: PRP (Properties)
        (unclaimed sequence; sequences of blood-coagulation factor VIIa-binding
        peptides)
AB
     This invention provides sequences of 6 blood-coagulation factor
     VIIa-binding peptides. This invention provides novel compds. which
     prevent or block a FVIIa mediated or associated process or event such as the
     catalytic conversion of FX to FXa , FVII to FVIIa or FIX to FIXa.
     particular aspects, the compds. of the invention bind Factor VIIa (FVIIa
     ), its zymogen Factor VII (FVII). The invention also provides
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685513-39-7P

RL: BPN (Biosynthetic preparation); PRP (Properties); THU (Therapeutic

(factor VIIa-binding anticoagulant peptide sequence; sequences of

use); BIOL (Biological study); PREP (Preparation); USES (Uses)

685513-40-0P

IT

358740-54-2P

685513-41-1P

685512-19-0P

685513-42-2P

pharmaceutical compns. comprising the novel compds. as well as their use in diagnostic, therapeutic, and prophylactic methods.

- L11 ANSWER 2 OF 4 CAPLUS COPYRIGHT 2005 ACS on STN
- AN 2003:443545 CAPLUS
- DN 139:159700
- TI Engineering Exosite Peptides for Complete Inhibition of Factor VIIa Using a Protease Switch with Substrate Phage
- AU Maun, Henry R.; Eigenbrot, Charles; Lazarus, Robert A.
- CS Department of Protein Engineering, Genentech, Inc., South San Francisco, CA, 94080, USA
- SO Journal of Biological Chemistry (2003), 278(24), 21823-21830 CODEN: JBCHA3; ISSN: 0021-9258
- PB American Society for Biochemistry and Molecular Biology
- DT Journal
- LA English
- IT 575431-91-3P, A 183X

RL: PAC (Pharmacological activity); PNU (Preparation, unclassified); THU (Therapeutic use); BIOL (Biological study); PREP (Preparation); USES (Uses)

(engineering exosite peptides for complete inhibition of factor VIIa using a protease switch with substrate phage)

AB Limitations of current anticoagulant therapies have led us to develop two distinct classes of exosite peptide inhibitors for the initiator of the clotting process, the tissue factor-factor VIIa (TF·FVIIa) complex (Roberge, M., Santell, L., Dennis, M. S., Eigenbrot, C., Dwyer, M. A., and Lazarus, R. A. (2001) Biochem. 40, 9522-9531). Although both peptide classes are potent and selective inhibitors of TF·FVIIa, neither showed 100% inhibition at saturating concns. Crystal structures of these peptides in complex with the FVII/FVIIa protease domain revealed their distinct binding sites and close proximity to the active site. The favorable orientation of the 15-mer A-site peptide A-183 (EEWEVLCWTWETCER) suggested that a C-terminal extension into the FVIIa active site could yield a chimeric inhibitor that was not only potent and selective but complete as well. A novel two-step "protease switch" approach using substrate phage display was developed by first binding all phage containing A-183 and C-terminal extension libraries to immobilized and inactive FVIIa. Upon altering pH and adding TF to switch on FVIIa enzymic activity, only those phage released by proteolytic cleavage within the extension were propagated. This process selected for both preferred sequence and length in the extension, leading to a 27-mer peptide A-183X (EEWEVLCWTWETCERGEGVEEELWEWR) with a C-terminal 12-mer extension containing an Arg in the P1 position. A-183X was a more potent and complete inhibitor of FX activation, having a maximal extent of inhibition of .apprx.99% with an IC50 of 230 pM vs. A-183 which maximally inhibited to 74% with an IC50 of 1.5 nM. A-183X also had a maximal prolongation of the prothrombin time of 7.6- vs. 1.9-fold for A-183, making it a more effective anticoagulant.

RE.CNT 44 THERE ARE 44 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT

- L11 ANSWER 3 OF 4 CAPLUS COPYRIGHT 2005 ACS on STN
- AN 2001:514493 CAPLUS
- DN 135:223287
- TI A novel exosite on coagulation factor VIIa and its molecular interactions with a new class of peptide inhibitors
- AU Roberge, Martin; Santell, Lydia; Dennis, Mark S.; Eigenbrot, Charles; Dwyer, Mary A.; Lazarus, Robert A.
- CS Department of Protein Engineering, Genentech Inc., South San Francisco, CA, 94080, USA
- SO Biochemistry (2001), 40(32), 9522-9531 CODEN: BICHAW; ISSN: 0006-2960
- PB American Chemical Society
- DT Journal
- LA English

ΙT 319927-97-4 325722-51-8 325722-64-3 358740-54-2 358740-54-2D, biotinylated derivs. RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); BIOL (Biological study) (protease domain exosite on coagulation factor VIIa and mol. interactions with A-series peptide inhibitors) A new inhibitory peptide binding exosite on the protease domain of AB coagulation Factor VIIa (FVIIa) has been identified. A novel series of peptide inhibitors of FVIIa, termed the "A-series" peptides, identified from peptide phage libraries and exemplified by peptide A-183, specifically bind at a site that is distinct from both the active site and the exosite of another recently described peptide inhibitor of FVIIa, E-76. Peptide A-183 prolonged TF-dependent clotting in human, but not rabbit plasma. Thus, a panel of human FVIIa mutants, containing 70 of the 76 rabbit sequence differences in the protease domain, localized the binding site to residues in the 60s loop and the C-terminus. The location of the exosite was refined by a series of FVIIa alanine mutants, which showed that proximal residues Trp 61 and Leu 251 were critical for binding. Kinetic and equilibrium binding consts. for zymogen FVII, FVIIa and TF FVIIa were determined using immobilized N-terminal biotinylated A-183 by surface plasmon resonance. No peptide binding to nine other human serine proteases was observed Key residues on the peptide were determined from binding to FVIIa and inhibition of FX activation using a series of alanine mutants of A-183 fused to the Z domain of protein A. Anal. of the mutagenesis data is presented in the context of a crystal structure of A-183 in complex with a version of zymogen FVII. The shape and proximity of this exosite to the active site may lend itself towards the design of new anticoaqulants that inhibit FVIIa. RE.CNT 47 THERE ARE 47 CITED REFERENCES AVAILABLE FOR THIS RECORD . ALL CITATIONS AVAILABLE IN THE RE FORMAT L11 ANSWER 4 OF 4 CAPLUS COPYRIGHT 2005 ACS on STN AN 2001:496925 CAPLUS DN 135:221051 TISelection and characterization of a new class of peptide exosite inhibitors of coaqulation factor VIIa ΑU Dennis, Mark S.; Roberge, Martin; Quan, Cliff; Lazarus, Robert A. CS Departments of Protein Engineering and Bioorganic Chemistry, Genentech Inc., South San Francisco, CA, 94080, USA SO Biochemistry (2001), 40(32), 9513-9521 CODEN: BICHAW; ISSN: 0006-2960 PB American Chemical Society DT Journal T.A English IT 325722-64-3 358740-54-2 359635-57-7 325722-51-8 359635-58-8 359635-59-9 359635-60-2 359635-61-3 359635-62-4 359635-63-5 359635-64-6 359635-65-7 359635-66-8 359635-67-9 RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); PRP (Properties); BIOL (Biological study) (selection and characterization of peptide exosite inhibitors of coagulation factor VIIa) A new series of peptide inhibitors of human Factor VIIa (FVIIa) has been AΒ identified and affinity matured from naive and partially randomized

An ew series of peptide inhibitors of human Factor VIIa (FVIIa) has been identified and affinity matured from naive and partially randomized peptide phage libraries selected against the immobilized tissue factor Factor VIIa (TF·FVIIa) complex. These "A-series" peptides contain a single disulfide bond and a 13-residue minimal core required for maximal affinity. They are exemplified by peptide A-183 (EEWEVLCWTWETCER), which binds at a newly identified exosite on the FVIIa protease domain, described in the accompanying report [Roberge, M., Santell, L., Dennis, M. S., Eigenbrot, C., Dwyer, M. A., and Lazarus, R. A. (2001) Biochem. 40, XXXXXX-XXXXX]. A-183 was obtained from a trypsin digest of A-100-Z, a recombinant protein comprising A-183 and the Z domain

of protein A. Surprisingly, A-183 was a very potent inhibitor of TF·FVIIa, inhibiting activation of Factor X (FX) and Factor IX and amidolytic activity of Chromozym t-PA with IC50 values of 1.6 \pm 1.2, 3.5 \pm 0.3, and 8.5 \pm 3.5 nM, resp. Kinetic anal. revealed that A-183 was a partial (hyperbolic) mixed-type inhibitor of FX activation having a Ki of 200 pM as well as a partial competitive inhibitor of amidolytic activity. The A-series peptides were also specific and potent inhibitors of TF-dependent clotting as measured in a prothrombin time (PT) clotting assay and had no effect on the TF-independent activated partial thromboplastin time. At saturating concns. of peptide, the maximal extent by which A-183 and A-100-Z inhibited the rate of FX activation was 78 \pm 3 and 89 + 6%, resp. The degree of inhibition of the rate of FX activation correlated with a maximum fold prolongation in the PT assay of 1.8-fold for A-183 and 3.3-fold for A-100-Z. The A-series peptides represent a new class of peptide exosite inhibitors that are capable of attenuating, rather than completely inhibiting, the activity of TF FVIIa, potentially leading to anticoagulants with an increased therapeutic window.

RE.CNT 41 THERE ARE 41 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT

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Crossover limits have been increased. See HELP CROSSOVER for details.

Experimental and calculated property data are now available. For more information enter HELP PROP at an arrow prompt in the file or refer to the file summary sheet on the web at: http://www.cas.org/ONLINE/DBSS/registryss.html

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COST IN U.S. DOLLARS

ENTRY SESSION 211.39 211.60

FULL ESTIMATED COST

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L11 ANSWER 1 OF 4 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 1

AN 2004:372929 CAPLUS

DN 140:395489

TI Sequences of blood-coagulation factor VIIa-binding peptides

IN Lazarus, Robert A.; Maun, Henry R.

PA Genentech, Inc., USA

SO U.S. Pat. Appl. Publ., 102 pp.

CODEN: USXXCO

DT Patent

LA English

FAN.CNT 1

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
PI US 2004087767	A1	20040506	US 2003-356257	20030130
PRAI US 2002-355420P	P	20020206		

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(factor VIIa-binding anticoagulant peptide sequence; sequences of
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     RL: PRP (Properties)
        (unclaimed sequence; sequences of blood-coagulation factor VIIa-binding
        peptides)
AB
     This invention provides sequences of 6 blood-coagulation factor
     VIIa-binding peptides. This invention provides novel compds. which
     prevent or block a FVIIa mediated or associated process or event such as the
     catalytic conversion of FX to FXa , FVII to FVIIa or FIX to FIXa.
     particular aspects, the compds. of the invention bind Factor VIIa (FVIIa
     ), its zymogen Factor VII (FVII). The invention also provides
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685513-39-7P

RL: BPN (Biosynthetic preparation); PRP (Properties); THU (Therapeutic

use); BIOL (Biological study); PREP (Preparation); USES (Uses)

685513-40-0P

685512-19-0P

685513-42-2P

IT

358740-54-2P

685513-41-1P

pharmaceutical compns. comprising the novel compds. as well as their use in diagnostic, therapeutic, and prophylactic methods.

- L11 ANSWER 2 OF 4 CAPLUS COPYRIGHT 2005 ACS on STN
- AN 2003:443545 CAPLUS
- DN 139:159700
- TI Engineering Exosite Peptides for Complete Inhibition of Factor VIIa Using a Protease Switch with Substrate Phage
- AU Maun, Henry R.; Eigenbrot, Charles; Lazarus, Robert A.
- CS Department of Protein Engineering, Genentech, Inc., South San Francisco, CA, 94080, USA
- SO Journal of Biological Chemistry (2003), 278(24), 21823-21830 CODEN: JBCHA3; ISSN: 0021-9258
- PB American Society for Biochemistry and Molecular Biology
- DT Journal
- LA English
- IT 575431-91-3P, A 183X

RL: PAC (Pharmacological activity); PNU (Preparation, unclassified); THU (Therapeutic use); BIOL (Biological study); PREP (Preparation); USES (Uses)

(engineering exosite peptides for complete inhibition of factor VIIa using a protease switch with substrate phage)

AΒ Limitations of current anticoagulant therapies have led us to develop two distinct classes of exosite peptide inhibitors for the initiator of the clotting process, the tissue factor-factor VIIa (TF·FVIIa) complex (Roberge, M., Santell, L., Dennis, M. S., Eigenbrot, C., Dwyer, M. A., and Lazarus, R. A. (2001) Biochem. 40, 9522-9531). Although both peptide classes are potent and selective inhibitors of TF·FVIIa, neither showed 100% inhibition at saturating concns. Crystal structures of these peptides in complex with the FVII/FVIIa protease domain revealed their distinct binding sites and close proximity to the active site. favorable orientation of the 15-mer A-site peptide A-183 (EEWEVLCWTWETCER) suggested that a C-terminal extension into the FVIIa active site could yield a chimeric inhibitor that was not only potent and selective but complete as well. A novel two-step "protease switch" approach using substrate phage display was developed by first binding all phage containing A-183 and C-terminal extension libraries to immobilized and inactive FVIIa. Upon altering pH and adding TF to switch on FVIIa enzymic activity, only those phage released by proteolytic cleavage within the extension were propagated. This process selected for both preferred sequence and length in the extension, leading to a 27-mer peptide A-183X (EEWEVLCWTWETCERGEGVEEELWEWR) with a C-terminal 12-mer extension containing an Arg in the P1 position. A-183X was a more potent and complete inhibitor of FX activation, having a maximal extent of inhibition of .apprx.99% with an IC50 of 230 pM vs. A-183 which maximally inhibited to 74% with an IC50 of 1.5 nM. A-183X also had a maximal prolongation of the prothrombin time of 7.6- vs. 1.9-fold for A-183, making it a more effective anticoagulant.

RE.CNT 44 THERE ARE 44 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT

- L11 ANSWER 3 OF 4 CAPLUS COPYRIGHT 2005 ACS on STN
- AN 2001:514493 CAPLUS
- DN 135:223287
- TI A novel exosite on coagulation factor VIIa and its molecular interactions with a new class of peptide inhibitors
- AU Roberge, Martin; Santell, Lydia; Dennis, Mark S.; Eigenbrot, Charles; Dwyer, Mary A.; Lazarus, Robert A.
- CS Department of Protein Engineering, Genentech Inc., South San Francisco, CA, 94080, USA
- SO Biochemistry (2001), 40(32), 9522-9531 CODEN: BICHAW; ISSN: 0006-2960
- PB American Chemical Society
- DT Journal
- LA English

IT 319927-97-4 325722-51-8 325722-64-3 358740-54-2 358740-54-2D, biotinylated derivs. RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); BIOL (Biological study) (protease domain exosite on coagulation factor VIIa and mol. interactions with A-series peptide inhibitors) A new inhibitory peptide binding exosite on the protease domain of AB coagulation Factor VIIa (FVIIa) has been identified. A novel series of peptide inhibitors of FVIIa, termed the "A-series" peptides, identified from peptide phage libraries and exemplified by peptide A-183, specifically bind at a site that is distinct from both the active site and the exosite of another recently described peptide inhibitor of FVIIa, E-76. Peptide A-183 prolonged TF-dependent clotting in human, but not rabbit plasma. Thus, a panel of human FVIIa mutants, containing 70 of the 76 rabbit sequence differences in the protease domain, localized the binding site to residues in the 60s loop and the C-terminus. The location of the exosite was refined by a series of FVIIa alanine mutants, which showed that proximal residues Trp 61 and Leu 251 were critical for binding. and equilibrium binding consts. for zymogen FVII, FVIIa and TF·FVIIa were determined using immobilized N-terminal biotinylated A-183 by surface plasmon resonance. No peptide binding to nine other human serine proteases was observed Key residues on the peptide were determined from binding to FVIIa and inhibition of FX activation using a series of alanine mutants of A-183 fused to the Z domain of protein A. Anal. of the mutagenesis data is presented in the context of a crystal structure of A-183 in complex with a version of zymogen FVII. The shape and proximity of this exosite to the active site may lend itself towards the design of new anticoagulants that inhibit FVIIa. RE.CNT 47 THERE ARE 47 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT L11 ANSWER 4 OF 4 CAPLUS COPYRIGHT 2005 ACS on STN ΑN 2001:496925 CAPLUS DN 135:221051 TI Selection and characterization of a new class of peptide exosite inhibitors of coagulation factor VIIa ΑU Dennis, Mark S.; Roberge, Martin; Quan, Cliff; Lazarus, Robert A. CS Departments of Protein Engineering and Bioorganic Chemistry, Genentech Inc., South San Francisco, CA, 94080, USA SO Biochemistry (2001), 40(32), 9513-9521 CODEN: BICHAW; ISSN: 0006-2960 PB American Chemical Society DT Journal LA English 325722-64-3 358740-54-2 359635-57-7 IT 325722-51-8 359635-58-8 359635-59-9 359635-60-2 359635-63-5 359635-61-3 359635-62-4 359635-64-6 359635-65-7 359635-66-8 359635-67-9 RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); PRP (Properties); BIOL (Biological study) (selection and characterization of peptide exosite inhibitors of coagulation factor VIIa)

AB A new series of peptide inhibitors of human Factor VIIa (FVIIa) has been identified and affinity matured from naive and partially randomized peptide phage libraries selected against the immobilized tissue factor Factor VIIa (TF·FVIIa) complex. These "A-series" peptides contain a single disulfide bond and a 13-residue minimal core required for maximal affinity. They are exemplified by peptide A-183 (EEWEVLCWTWETCER), which binds at a newly identified exosite on the FVIIa protease domain, described in the accompanying report [Roberge, M., Santell, L., Dennis, M. S., Eigenbrot, C., Dwyer, M. A., and Lazarus, R. A. (2001) Biochem. 40, XXXXXX-XXXXX]. A-183 was obtained from a trypsin digest of A-100-Z, a recombinant protein comprising A-183 and the Z domain

of protein A. Surprisingly, A-183 was a very potent inhibitor of TF·FVIIa, inhibiting activation of Factor X (FX) and Factor IX and amidolytic activity of Chromozym t-PA with IC50 values of 1.6 \pm 1.2, 3.5 \pm 0.3, and 8.5 \pm 3.5 nM, resp. Kinetic anal. revealed that A-183 was a partial (hyperbolic) mixed-type inhibitor of FX activation having a Ki of 200 pM as well as a partial competitive inhibitor of amidolytic activity. The A-series peptides were also specific and potent inhibitors of TF-dependent clotting as measured in a prothrombin time (PT) clotting assay and had no effect on the TF-independent activated partial thromboplastin time. At saturating concns. of peptide, the maximal extent by which A-183 and A-100-Z inhibited the rate of FX activation was 78 \pm 3 and 89 ± 6%, resp. The degree of inhibition of the rate of FX activation correlated with a maximum fold prolongation in the PT assay of 1.8-fold for A-183 and 3.3-fold for A-100-Z. The A-series peptides represent a new class of peptide exosite inhibitors that are capable of attenuating, rather than completely inhibiting, the activity of TF·FVIIa, potentially leading to anticoagulants with an increased therapeutic window.

RE.CNT 41 THERE ARE 41 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT

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L13 ANSWER 1 OF 2 CAPLUS COPYRIGHT 2005 ACS on STN
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     (Biological study); PROC (Process)
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(factor VIIa antagonists for diagnostic or therapeutic use)

=> d l13 hit bib 1-2

L13 ANSWER 1 OF 2 CAPLUS COPYRIGHT 2005 ACS on STN

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RL: BAC (Biological activity or effector, except adverse); BPR (Biological
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     (Biological study); PROC (Process)
        (factor VIIa antagonists for diagnostic or therapeutic use)
AN
     2001:115174 CAPLUS
DN
     134:168300
     Factor VIIa antagonists for diagnostic or therapeutic use
TI
IN
     Dennis, Mark S.
PΑ
     Genentech, Inc., USA
SO
     PCT Int. Appl., 80 pp.
     CODEN: PIXXD2
DT
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LA
     English
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L13 ANSWER 2 OF 2 USPATFULL on STN
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    325722-42-7
        (factor VIIa antagonists for diagnostic or therapeutic use)
       2004:101678 USPATFULL
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       FVIIa antagonists
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PA
       Genentech, Inc., South San Francisco, CA (U.S. corporation)
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DT
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FS
       APPLICATION
       GENENTECH, INC., 1 DNA WAY, SOUTH SAN FRANCISCO, CA, 94080
LREP
CLMN
       Number of Claims: 31
ECL
       Exemplary Claim: 1
DRWN
       4 Drawing Page(s)
LN.CNT 2987
CAS INDEXING IS AVAILABLE FOR THIS PATENT.
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IT

325722-42-7

COST IN U.S. DOLLARS SINCE FILE TOTAL

ENTRY SESSION FULL ESTIMATED COST 27.77 239.37

DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS) SINCE FILE TOTAL

ENTRY SESSION CA SUBSCRIBER PRICE

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STRUCTURE FILE UPDATES: 3 JAN 2005 HIGHEST RN 807382-78-1 DICTIONARY FILE UPDATES: 3 JAN 2005 HIGHEST RN 807382-78-1

TSCA INFORMATION NOW CURRENT THROUGH MAY 21, 2004

Please note that search-term pricing does apply when conducting SmartSELECT searches.

Crossover limits have been increased. See HELP CROSSOVER for details.

Experimental and calculated property data are now available. For more information enter HELP PROP at an arrow prompt in the file or refer to the file summary sheet on the web at: http://www.cas.org/ONLINE/DBSS/registryss.html

=> s [wfl]e[lv][limv]c[wflm]twetce[rklw]/sqsp 24 [WFL] E [LV] [LIMV] C [WFLM] TWETCE [RKLW] / SQSP

-2.92

-2.92

=> s [wfla]e[via]lc[wflma]twetcer/sqsp 22 [WFLA] E [VIA] LC [WFLMA] TWETCER/SQSP

=> FIL CAPLUS BIOSIS MEDLINE PCTFULL USPATFULL EUROPATFULL JAPIO SCISEARCH EMBASE COST IN U.S. DOLLARS SINCE FILE TOTAL

ENTRY SESSION FULL ESTIMATED COST 58.87 298.24

DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS) SINCE FILE TOTAL ENTRY SESSION CA SUBSCRIBER PRICE 0.00 -2.92

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FILE 'JAPIO' ENTERED AT 15:17:14 ON 04 JAN 2005 COPYRIGHT (C) 2005 Japanese Patent Office (JPO) - JAPIO

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=> FIL REGISTRY

COST IN U.S. DOLLARS

SINCE FILE TOTAL
ENTRY SESSION

FULL ESTIMATED COST

DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS)

SINCE FILE TOTAL
ENTRY SESSION
SESSION

0.00

-2.92

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Property values tagged with IC are from the ZIC/VINITI data file provided by InfoChem.

STRUCTURE FILE UPDATES: 3 JAN 2005 HIGHEST RN 807382-78-1 DICTIONARY FILE UPDATES: 3 JAN 2005 HIGHEST RN 807382-78-1

TSCA INFORMATION NOW CURRENT THROUGH MAY 21, 2004

Please note that search-term pricing does apply when conducting SmartSELECT searches.

Crossover limits have been increased. See HELP CROSSOVER for details.

Experimental and calculated property data are now available. For more information enter HELP PROP at an arrow prompt in the file or refer to the file summary sheet on the web at: http://www.cas.org/ONLINE/DBSS/registryss.html

=> d hist

CA SUBSCRIBER PRICE

(FILE 'HOME' ENTERED AT 14:50:12 ON 04 JAN 2005)

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L10 5 S L9

L11 4 DUP REM L10 (1 DUPLICATE REMOVED)

L12 7 S L2

L13 2 S L12 NOT L10

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24 S [WFL] E [LV] [LIMV] C [WFLM] TWETCE [RKLW] /SQSP

L15 22 S [WFLA] E [VIA] LC [WFLMA] TWETCER/SQSP

FILE 'CAPLUS, BIOSIS, MEDLINE, PCTFULL, USPATFULL, JAPIO, SCISEARCH, EMBASE' ENTERED AT 15:17:14 ON 04 JAN 2005

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=> s l14 or l15

L14

L16 28 L14 OR L15

=> FIL CAPLUS BIOSIS MEDLINE PCTFULL USPATFULL EUROPATFULL JAPIO SCISEARCH EMBASE USPAT2 EUROPATFULL

COST IN U.S. DOLLARS

SINCE FILE TOTAL ENTRY SESSION

FULL ESTIMATED COST

0.43 307.70

DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS)

SINCE FILE TOTAL

CA SUBSCRIBER PRICE

ENTRY SESSION 0.00 -2.92

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CA INDEXING COPYRIGHT (C) 2005 AMERICAN CHEMICAL SOCIETY (ACS).

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L17 7 L14

^{=&}gt; s 114

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L23
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DN
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ΤI
     Sequences of blood-coagulation factor VIIa-binding peptides
     Lazarus, Robert A.; Maun, Henry R.
IN
     Genentech, Inc., USA
PA
     U.S. Pat. Appl. Publ., 102 pp.
SO
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    English
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     RL: PRP (Properties)
        (unclaimed sequence; sequences of blood-coagulation factor VIIa-binding
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L23 ANSWER 2 OF 6 USPATFULL on STN
      2004:101678 USPATFULL
      FVIIa antagonists
      Dennis, Mark S., San Carlos, CA/ UNITED STATES
      Genentech, Inc., South San Krancisco, CA (U.S. corporation)
                              20/14/22
      US 2004077547
                         A1
                              20030 (10)
      US 2003-639076
                         Α1
      Continuation of Ser. No. U$ 200Q-632429, filed on 4 Aug 2000, PENDING
                          19990800 (60)
      US 1999-147627P
      US 1999-150315P
                          19990823 (60)
      Utility
      APPLICATION
      GENENTECH, INC., 1 DNA WAY, SOUTH SAN FRANCISCO, CA, 94080
      Number of Claims: 31
      Exemplary Claim: 1
       4 Drawing Page(s)
LN.CNT 2987
CAS INDEXING IS AVAILABLE FOR THIS PATENT.
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RLI

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LREP CLMN

ECL

DRWN

PRAI

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IT 325722-42-7
        (factor VIIa antagonists for diagnostic or therapeutic use)
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AN
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     139:159700
DN
     Engineering Exosite Peptides for Complete Inhibition of Factor VIIa Using
TI
     a Protease Switch with Substrate Phage
ΑU
     Maun, Henry R.; Eigenbrot, Charles; Lazarus, Robert A.
     Department of Protein Engineering, Genentech, Inc., South San Francisco,
CS
     CA, 94080, USA
                                     (2003), 278(24), 21823-21830
so
     Journal of Biological Chemistry
     CODEN: JBCHA3; ISSN: 0021-9258
PB
     American Society for Biochemistry and Molecular Biology
DT
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LA
     English
RE.CNT 44
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     RL: PAC (Pharmacological activity); PNU (Preparation, unclassified); THU
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        using a protease switch with substrate phage)
L23
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     134:168300
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IN
     Dennis, Mark S.
     Genentech, Inc., USA
PΑ
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     PCT Int. Appl., 80 pp.
     CODEN: PIXXD2
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     English
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                                DATE
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IT
     325722-42-7
     RL: BAC (Biological activity or effector, except adverse); BPR (Biological
     process); BSU (Biological study, unclassified); PRP (Properties); BIOL
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(factor VIIa antagonists for diagnostic or therapeutic use) L23 ANSWER 5 OF 6 CAPLUS COPYRIGHT 2005 ACS on STN 2001:514493 CAPLUS ANDN 135:223287 A novel exosite on coagulation factor VIIa and its molecular interactions TI with a new class of peptide inhibitors AU Roberge, Martin; Santell, Lydia; Dennis, Mark S.; Eigenbrot, Charles; Dwyer, Mary A.; Lafarus, Robert A. Department of Provein Engineering, Genentech Inc., South San Francisco, CS CA, 94080, USA

Biochemistry (2001), 40(32), 9522-9531

CODEN: BICHAW; VSN: 0006-2960 SO PB American Chemical Society DTJournal English LA RE.CNT 47 THERE ARE 47 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT IT 319927-97-4 325722-51-8 325722-64-3 **358740-54-2** 358740-54-2D, biotinylated derivs. RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); BIOL (Biological study) (protease domain exosite on coagulation factor VIIa and mol. interactions with A-series peptide inhibitors) ANSWER 6 OF 6 CAPLUS COPYRIGHT 2005 ACS on STN L23 AN 2001:496925 CAPLUS DN 135:221051 Selection and characterization of a new class of peptide exosite ΤI inhibitors of coagulation factor VIIa Dennis, Mark S.; Roberge, Martin; Quan, Cliff; Lazarus, Robert A. Departments of Protein Engineering and Bioorganic Chemistry, Genentech ΔII CS Inc., South San Ffancisco, CA, 94080, USA Biochemistry (2001), 40(32), 9513-9521 CODEN: BICHAW; USAN: 0006-2960 SO American Chemical Society PB DT Journal LAEnglish RE.CNT 41 THERE ARE 41 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT IT 325722-51-8 325722-64-3 **358740-54-2 359635-57-7** 359635-58-8 359635-59-9 359635-60-2 359635-61-3 359635-62-4 359635-63-5 359635-64-6 359635-67-9 359635-65-7 359635-66-8 RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); PRP (Properties); BIOL (Biological study) (selection and characterization of peptide exosite inhibitors of coagulation factor VIIa) => FIL REGISTRY COST IN U.S. DOLLARS SINCE FILE TOTAL ENTRY SESSION FULL ESTIMATED COST 36.80 344.50 DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS) SINCE FILE TOTAL ENTRY SESSION CA SUBSCRIBER PRICE 0.00 -2.92

(Biological study); PROC (Process)

FILE 'REGISTRY' ENTERED AT 15:21:32 ON 04 JAN 2005

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STRUCTURE FILE UPDATES: 3 JAN 2005 HIGHEST RN 807382-78-1 DICTIONARY FILE UPDATES: 3 JAN 2005 HIGHEST RN 807382-78-1

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Crossover limits have been increased. See HELP CROSSOVER for details.

Experimental and calculated property data are now available. For more information enter HELP PROP at an arrow prompt in the file or refer to the file summary sheet on the web at: http://www.cas.org/ONLINE/DBSS/registryss.html

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=> FIL CAPLUS BIOSIS MEDLINE PCTFULL USPATFULL EUROPATFULL JAPIO SCISEARCH EMBASE COST IN U.S. DOLLARS SINCE FILE TOTAL

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PΒ Kazusa DNA Research Institute DT Journal LΑ English THERE ARE 30 CITED REFERENCES AVAILABLE FOR THIS RECORD RE.CNT 30 ALL CITATIONS AVAILABLE IN THE RE FORMAT => d 130 bib hit L30 ANSWER 1 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN 2004:1012134 CAPLUS AN DN 141:421056 Expressed sequence tags and encoded human proteins TITN Edwards, Jean-Baptiste Dumas Milne; Duclert, Aymeric; Giordano, Jean-Yves PΑ Genset S.A., Fr. U.S., 72 pp., Cont.-in-part of Appl. No. PCT/IB99/00712. SO CODEN: USXXAM DT Patent LA English FAN.CNT 2 PATENT NO. KIND DATE APPLICATION NO. DATE ΡI US 6822072 В1 20041123 US 1999-471276 19991221 WO 9953051 A2 19991021 WO 1999-IB712 19990409 <--A3 WO 9953051 20000406 W: AU, CA, JP, US RW: AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE PRAI US 1998-57719 B2 19980409 · B2 US 1998-69047 19980428 WO 1999-IB712 A2 19990409 RE.CNT 22 THERE ARE 22 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT PATENT NO. KIND DATE APPLICATION NO. DATE _ _ _ _ _____ -----_____ US 1999-471276 19991221 PΙ US 6822072 В1 20041123 WO 9953051 WO 1999-IB712 19990409 <--A2 19991021 WO 9953051 Α3 20000406 W: AU, CA, JP, US RW: AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE IT 246537-99-5 220593-06-6 225514-15-8 246537-95-1 246538-06-7 246538-08-9 246538-10-3 246538-24-9 246538-07-8 246538-09-0 246538-74-9 246538-77-2 246877-53-2 246877-54-3 246877-55-4 246877-58-7 **246877-59-8** 246877-56-5 246877-57-6 246877-64-5 246877-60-1 246877-61-2 246877-62-3 246877-63-4 246877-65-6 246877-66-7 246877-67-8 246877-68-9 246877-69-0 246877-70-3 246877-71-4 246877-72-5 246877-73-6 246877-74-7 246877-75-8 246877-76-9 246877-77-0 246877-78-1 246877-79-2 246877-80-5 246877-81-6 246877-82-7 246877-83-8 246877-84-9 246877-85-0 246877-86-1 246877-87-2 246877-88-3 246877-89-4 246877-90-7 246877-91-8 246877-92-9 246877-93-0 246877-94-1 246877-95-2 246877-96-3 246877-97-4 246877-98-5 246877-99-6 246878-00-2 246878-01-3 246878-02-4 246878-03-5 246878-04-6 246878-05-7 246878-06-8 246878-07-9 246878-08-0 246878-09-1 246878-10-4 246878-11-5 246878-12-6 246878-13-7 246878-14-8 246878-15-9 246878-16-0 246878-17-1 246878-18-2 246878-19-3 246878-20-6 246878-21-7 246878-22-8 246878-23-9 246878-24-0 246878-25-1 246878-26-2 246878-27-3 246878-28-4 246878-29-5 246878-30-8 246878-31-9 246878-32-0 246878-33-1 246878-34-2

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    De Waal, Malefyt Rene; Flickensher, Helmut; Fleckenstein, Bernhard;
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    Schering Corp., USA
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L30
     1999:723169 CAPLUS
AN
DN
     131:348189
ТT
     Human receptor (REC) polypeptides and polynucleotides, sequences, and
     biological and therapeutic uses thereof
TN
     Hillman, Jennifer L.; Bandman, Olga; Tang, Y. Tom; Yue, Henry; Lal,
     Preeti; Corley, Neil C.; Guegler, Karl J.; Patterson, Chandra
PΑ
     Incyte Pharmaceuticals, Inc., USA
SO
     PCT Int. Appl., 94 pp.
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     1999:673017 CAPLUS
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ΤI
     5'-Expressed sequence tags for secreted proteins identified from human
IN
     Dumas Milne Edwards, Jean-Baptiste; Duclert, Aymeric; Giordano, Jean-Yves
PA
     Genset S. A., Fr.
SO
     PCT Int. Appl., 837 pp.
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AN
     1999:614249 CAPLUS
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     131:252536
     Assays for screening compounds which interact with cation channel
ΤI
     proteins, mutant prokaryotic cation channel proteins, and uses thereof
TN
     MacKinnon, Roderick
PΑ
     The Rockefeller University, USA
so
     PCT Int. Appl., 165 pp.
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AN
     1999:614169 CAPLUS
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     Human and murine G protein-coupled heptahelical receptor D6 and its cDNA
TI
     sequences and therapeutic uses
     Graham, Gerard J.; Benjamin, Nibbs Robert J.; Gonzalo, Jose-Angel;
IN
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Gutierrez-Ramos, Jose-Carlos

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SO
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             THERE ARE 4 CITED REFERENCES AVAILABLE FOR THIS RECORD
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    ANSWER 17 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
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    1999:576793 CAPLUS
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    131:195462
    Protein and cDNA sequences for human and mouse IL-9 Induced Calcium
TI
    Activated Chloride Channels (ICACC) and uses thereof in the treatment of
     atopic allergies, asthma, inflammatory bowel disease, and cystic fibrosis
IN
    Holroyd, Kenneth J.; Levitt, Roy C.; Maloy, W. Lee; Louahed, Jamila;
    McLane, Mike; Nicolaides, Nicholas C.; Zhou, Yuhong; Dong, Qu
PA
    Magainin Pharmaceuticals, Inc., USA
    PCT Int. Appl., 75 pp.
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Millennium Pharmaceuticals, Inc., USA; CRC Technology Limited

PA

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     ANSWER 18 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
AN
     1999:388051 CAPLUS
DN
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TI
     Phaseolus genes expressed during senescence and their promoters and the
     stage-specific expression of foreign genes
     Gepstein, Shimon; Hajuoje, Taleb; Rosner, Amalia
IN
PΑ
     Vitality Biotechnologies, Inc., USA
so
     PCT Int. Appl., 69 pp.
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     1999:48801 CAPLUS
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DN
     130:120476
TI
     Genes for Niemann-Pick type C disease
     Carstea, Eugene D.; Tagle, Danilo A.; Morris, Jill A.; Pentchev, Peter G.;
IN
     Pavan, William J.; Rosenfeld, Melissa A.; Loftus, Stacie K.; Gu, Jessie
     United States Dept. of Health and Human Services, USA
PΑ
SO
     PCT Int. Appl., 100 pp.
     CODEN: PIXXD2
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AN
     1999:802470 CAPLUS
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     132:45817
ΤI
     Cloning of gene for cytochrome bd type quinol oxidase from Brevibacterium
     lactofermentum
IN
     Sone, Nobufumi
PA
     Ajinomoto Co., Inc., Japan
SO
     Jpn. Kokai Tokkyo Koho, 19 pp.
     CODEN: JKXXAF
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2000050<u>.</u>
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PRAI JP 1998-164019
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    ANSWER 21 OF 224 USPATFULL on STN
L30
       Proteins involved in the synthesis and assembly of O-antigen in
TI
       Pseudomonas aeruginosa
IN
      Lam, Joseph S., Guelph, Canada
      Burrows, Lori, Guelph, Canada
       Charter, Deborah, Guelph, Canada
       de Kievit, Teresa, Guelph, Canada
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PΙ
      US 5994072
                             19991130
    ANSWER 22 OF 224 USPATFULL on STN
L30
      Rolling mill roll stand
TI
       Woodrow, Harold E., Northboro, MA, United States
IN
       Shore, T. Michael, Princeton, MA, United States
      US 5983694
PΙ
                             19991116
                                                                 <--
    ANSWER 23 OF 224 USPATFULL on STN
L30
      Fine magnetic particles containing useful proteins bound thereto,
TΙ
      process for producing the same, and use thereof
      Matsunaga, Tadashi, B-506, 2-40, Saiwai-cho, Funchi-shi, Tokyo, 183,
IN
       Japan
      Kamiya, Shinji, Tokyo, Japan
      Namba, Kenryo, Tokyo, Japan
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PI
      US 5958706
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      WO 9735964 19971002
    ANSWER 24 OF 224 USPATFULL on STN
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DNA encoding a 2-acyltransferases

TI

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- SO Biochemical and Biophysical Research Communications (1999), 261(2), 493-498
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       US 5810530
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     screening for ligands of the receptor
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     Tomalski, Michael D.; Gant, Daniel B.
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     CODEN: USXXAM
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     Stuiver, Maarten Hendrik; Custers, Jerome Hubertus Henricus Victor;
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B1 20041117
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A 20000327
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    NZ 334517
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    Knappe, Andrea; Fickenscher, Helmut; Fleckenstein, Bernard
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    CODEN: PIXXD2
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L30 ANSWER 48 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
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    jannaschii
    Bult, Carol J.; White, Owen R.; Smith, Hamilton O.; Woese, Carl R.;
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                     Venter, J. Craig
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L30 ANSWER 49 OF 224 USPATFULL on STN
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      DNA encoding 2-acyltransferases
IN
      Slabas, Antoni Ryszard, High Shincliffe, United Kingdom
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Interference blind type bolt

Brown, Adrian Paul, Shadforth, United Kingdom

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19981201

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     Li, Yi; Kirkness, Ewen F.
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     U.S., 29 pp.
     CODEN: USXXAM
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     US 5654172
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     Cascieri, Margaret A.; Linemeyer, David L.; Macneil, Douglas J.; Shiao,
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L30 ANSWER 72 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
     C14 sterol reductases of plants, nucleic acids encoding them, and
ΤI
     transgenic plants with altered C14 sterol reductase levels
     Jang, Jyan-Chyun; Sheen, Jen
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SO
     PCT Int. Appl., 71 pp.
     CODEN: PIXXD2
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L30 ANSWER 73 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
ΤI
    Proteins involved in the synthesis and assembly of the O-antigen of
     Pseudomonas aeruginosa and the genes encoding them
IN
     Lam, Joseph S.; Burrows, Lori; Charter, Deborah; De Kievit, Teresa
SO
     PCT Int. Appl., 194 pp.
     CODEN: PIXXD2
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    Preparation of proteins by expression of a fusion protein containing a
    membrane-binding protein of magnetic bacteria and use of the fusion
    protein
IN
    Matsunaga, Tadashi; Kamiya, Shinji; Namba, Kenryo
SO
    PCT Int. Appl., 70 pp.
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IN SO		s, Lori; Charter, Deborah; De Kievit, Teresa D.	
	PATENT NO. KIN		
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so	Journal of Biological C CODEN: JBCHA3; ISSN: 00	Chemistry (1997), 272(24), 15346-15350 021-9258	
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so		United Kingdom) (1997), 143(11),	
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L30· TI	Identification and char	JS COPYRIGHT 2005 ACS on STN racterization of a DNA region involved in the exp	
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- SO Neuron (1997), 19(5), 1061-1075 CODEN: NERNET; ISSN: 0896-6273
- L30 ANSWER 85 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
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- SO DNA and Cell Biology (1997), 16(9), 1023-1030 CODEN: DCEBE8; ISSN: 1044-5498
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- SO Journal of Biochemistry (Tokyo) (1997), 122(2), 438-452 CODEN: JOBIAO; ISSN: 0021-924X
- L30 ANSWER 89 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
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- SO DNA Research (1997), 4(2), 91-113, 169-178 CODEN: DARSE8; ISSN: 1340-2838
- L30 ANSWER 96 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
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- SO Gene (1997), 197(1/2), 47-64 CODEN: GENED6; ISSN: 0378-1119
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- SO Virus Research (1997), 47(1), 7-17 CODEN: VIREDF; ISSN: 0168-1702
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- IN Soderlund, David M.; Knipple, Douglas C.; Henderson, Joseph E.
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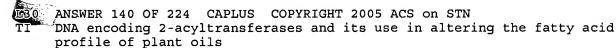
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- TI The genome organization of potato virus M RNA
- SO Journal of General Virology (1991), 72(1), 9-14 CODEN: JGVIAY; ISSN: 0022-1317
- L30 ANSWER 193 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN

- TI Identification and functional expression of a novel ligand binding subunit of the inhibitory glycine receptor
- SO Journal of Biological Chemistry (1990), 265(36), 22317-20 CODEN: JBCHA3; ISSN: 0021-9258
- L30 ANSWER 194 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Sequence of the chicken GABAA receptor γ2-subunit cDNA
- SO Nucleic Acids Research (1990), 18(23), 7157 CODEN: NARHAD; ISSN: 0305-1048
- L30 ANSWER 195 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Sulfate and thiosulfate transport in Escherichia coli K-12: nucleotide sequence and expression of the cysTWAM gene cluster
- SO Journal of Bacteriology (1990), 172(6), 3351-7 CODEN: JOBAAY; ISSN: 0021-9193
- L30 ANSWER 196 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Structural and functional characterization of the γ 1 subunit of GABAA/benzodiazepine receptors
- SO EMBO Journal (1990), 9(10), 3261-7 CODEN: EMJODG; ISSN: 0261-4189
- L30 ANSWER 197 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Functional characteristics and sites of gene expression of the $\alpha 1, \beta 1, \gamma 2$ -isoform of the rat GABAA receptor
- SO Journal of Neuroscience (1990), 10(7), 2330-7 CODEN: JNRSDS; ISSN: 0270-6474
- L30 ANSWER 198 OF 224 MEDLINE on STN
- TI Molecular analysis of the Escherichia coli K5 kps locus: identification and characterization of an inner-membrane capsular polysaccharide transport system.
- SO Molecular microbiology, **(1990 Nov)** 4 (11) 1863-9. Journal code: 8712028. ISSN: 0950-382X.
- L30 ANSWER 199 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 7
- TI The bex locus in encapsulated Haemophilus influenzae: a chromosomal region involved in capsule polysaccharide export
- SO Molecular Microbiology (1990), 4(11), 1853-62 CODEN: MOMIEE; ISSN: 0950-382X
- L30 ANSWER 200 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Cloning and expression of the 58 kd β subunit of the inhibitory glycine receptor
- SO Neuron (1990), 4(6), 963-70 CODEN: NERNET; ISSN: 0896-6273
- L30 ANSWER 201 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI A single amino acid exchange alters the pharmacology of neonatal rat glycine receptor subunit
- SO Neuron (1990), 5(6), 867-73 CODEN: NERNET; ISSN: 0896-6273
- L30 ANSWER 202 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Alpha subunit variants of the human glycine receptor: primary structures, functional expression and chromosomal localization of the corresponding genes
- SO EMBO Journal (1990), 9(3), 771-6 CODEN: EMJODG; ISSN: 0261-4189
- L30 ANSWER 203 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Nucleotide sequence and genome structure of the 3'-terminal region of potato virus M genomic RNA
- SO Molekulyarnaya Biologiya (Moscow) (1990), 24(2), 448-59

CODEN: MOBIBO; ISSN: 0026-8984

- L30 ANSWER 204 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Diversity of the Chlamydia trachomatis common plasmid in biovars with different pathogenicity
- SO Plasmid (1990), 23(2), 149-54 CODEN: PLSMDX; ISSN: 0147-619X
- L30 ANSWER 205 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI New mre genes mreC and mreD, responsible for formation of the rod shape of Escherichia coli cells
- SO Journal of Bacteriology (1989), 171(12), 6511-16 CODEN: JOBAAY; ISSN: 0021-9193
- L30 ANSWER 206 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Partial nucleotide sequence of potato virus M RNA shows similarities to potexviruses in gene arrangement and the encoded amino acid sequences
- SO Journal of General Virology (1989), 70(7), 1861-9 CODEN: JGVIAY; ISSN: 0022-1317
- L30 ANSWER 207 OF 224 CAPLUS COPYRIGHT 2005 ACS ON STN
- TI Importance of a novel GABAA receptor subunit for benzodiazepine pharmacology
- SO Nature (London, United Kingdom) (1989), 338(6216), 582-5 CODEN: NATUAS; ISSN: 0028-0836
- L30 ANSWER 208 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Two novel GABAA receptor subunits exist in distinct neuronal subpopulations
- SO Neuron (1989), 3(3), 327-37 CODEN: NERNET; ISSN: 0896-6273
- L30 ANSWER 209 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Analysis of the entire nucleotide sequence of the cryptic plasmid of Chlamydia trachomatis serovar L1. Evidence for involvement in DNA replication
- SO Nucleic Acids Research (1988), 16(9), 4053-67 CODEN: NARHAD; ISSN: 0305-1048
- L30 ANSWER 210 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Nucleotide sequence of the dmsABC operon encoding the anaerobic dimethylsulphoxide reductase of Escherichia coli
- SO Molecular Microbiology (1988), 2(6), 785-95 CODEN: MOMIEE; ISSN: 0950-382X
- L30 ANSWER 211 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Transient expression and sequence of the matrix (M1) gene of WSN influenza A virus in a vaccinia vector
- SO Virology (1988), 163(2), 618-21 CODEN: VIRLAX; ISSN: 0042-6822
- L30 ANSWER 212 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI The structure of a plasmid of Chlamydia trachomatis believed to be required for growth within mammalian cells
- SO Molecular Microbiology (1988), 2(4), 531-8 CODEN: MOMIEE; ISSN: 0950-382X
- L30 ANSWER 213 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Nucleotide sequence of RNA segment 7 and the predicted amino sequence of M1 and M2 proteins of FPV/Weybridge (H7N7) and WSN (H1N1) influenza viruses
- SO Virus Research (1988), 10(2-3), 263-71 CODEN: VIREDF; ISSN: 0168-1702

- L30 ANSWER 214 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Structure of Escherichia coli dnaC. Identification of a cysteine residue possibly involved in association with dnaB protein
- SO Journal of Biological Chemistry (1987), 262(22), 10475-80 CODEN: JBCHA3; ISSN: 0021-9258
- L30 ANSWER 215 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Sequence and regulatory responses of a ribosomal protein gene from the fission yeast Schizosaccharomyces pombe
- SO Nucleic Acids Research (1987), 15(4), 1477-92 CODEN: NARHAD; ISSN: 0305-1048
- L30 ANSWER 216 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Similarity between cell-cycle genes of budding yeast and fission yeast and the Notch gene of Drosophila
- SO Nature (London, United Kingdom) (1987), 329(6140), 651-4 CODEN: NATUAS; ISSN: 0028-0836
- L30 ANSWER 217 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI The strychnine-binding subunit of the glycine receptor shows homology with nicotinic acetylcholine receptors
- SO Nature (London, United Kingdom) (1987), 328(6127), 215-20 CODEN: NATUAS; ISSN: 0028-0836
- L30 ANSWER 218 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI The organization of the araBAD operon of Escherichia coli
- SO Gene (1986), 47(2-3), 231-44 CODEN: GENED6; ISSN: 0378-1119
- L30 ANSWER 219 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI The araBAD operon of Salmonella typhimurium LT2. I. Nucleotide sequence of araB and primary structure of its product, ribulokinase
- SO Gene (1985), 34(1), 111-22 CODEN: GENED6; ISSN: 0378-1119
- L30 ANSWER 220 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI The sequence of the gene for cytochrome c oxidase subunit I, a frameshift containing gene for cytochrome c oxidase subunit II and seven unassigned reading frames in Trypanosoma brucei mitochondrial maxi-circle DNA
- SO Nucleic Acids Research (1984), 12(19), 7327-44 CODEN: NARHAD; ISSN: 0305-1048
- L30 ANSWER 221 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Nucleotide sequence of an infectious molecularly cloned genome of ground squirrel hepatitis virus
- SO Journal of Virology (1984), 51(2), 367-75 CODEN: JOVIAM; ISSN: 0022-538X
- L30 ANSWER 222 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Somatic mutations of immunoglobulin variable genes are restricted to the rearranged V gene
- SO Science (Washington, DC, United States) (1983), 220(4602), 1179-81
 CODEN: SCIEAS; ISSN: 0036-8075
- L30 ANSWER 223 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Complete nucleotide sequences of cloned copies of the RNA genes coding for the hemagglutinin and matrix proteins of a human influenza virus
- SO Developments in Cell Biology (Amsterdam) (1981), 7 (Replication Negat. Strand Viruses), 241-9
 CODEN: DCBIDD; ISSN: 0165-2265
- L30 ANSWER 224 OF 224 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Cloning of influenza cDNA into M13: the sequence of the RNA segment

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encoding the A/PR/8/34 matrix protein
SO
     Nucleic Acids Research (1980), 8(9), 1965-74
     CODEN: NARHAD; ISSN: 0305-1048
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L4
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L5
L6
               5 S .{0,99}.WEVLCWTWETCER.{0,99}./SQSP
L7
               8 S . {0,99}.WEVLCWTWETCER/SQSP
L8
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L9
               9 S L6 OR L7 OR L8
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L10
L11
               4 DUP REM L10 (1 DUPLICATE REMOVED)
L12
              7 S L2
L13
              2 S L12 NOT L10
     FILE 'REGISTRY' ENTERED AT 15:12:05 ON 04 JAN 2005
L14
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L15
              22 S [WFLA] E [VIA] LC [WFLMA] TWETCER/SQSP
     FILE 'CAPLUS, BIOSIS, MEDLINE, PCTFULL, USPATFULL, JAPIO, SCISEARCH,
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     FILE 'REGISTRY' ENTERED AT 15:17:27 ON 04 JAN 2005
L16
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L17
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T.18
L19
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L20
T<sub>2</sub>1
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              7 S L21 AND L17
T<sub>2</sub>2
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L23
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L24
L25
         134568 S .. [WFLA] . [VIA] [LIMVA] [WFLMA] . [WFYM] .. [-P] [RKLWHM] .../SQSP
           2424 S .. [WFL] . [VI] [LIMV] [WFLM] . [W] .. [-P] [RKLW] .../SQSP
L26
           3807 S [WFL].[VI] [LIMVA] [WFYLM].[W]...[RKLW]/SQSP
L27
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            231 S L26 AND PY<=1999
L28
              0 S L28 AND DUP REM
L29
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224 DUP REM L28 (7 DUPLICATES REMOVED)

0 S L30 AND (FACTOR (W) VII)

1 S L30 AND (VII)

=> FIL REGISTRY
COST IN U.S. DOLLARS

L30 L31

L32

SINCE FILE TOTAL ENTRY SESSION

FULL ESTIMATED COST 470.73 932.54

DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS) SINCE FILE TOTAL

ENTRY

SESSION

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136536 SQL=18

L33 0 .. [WFL] . [VI] [LIMV] [WFLM] . [-P] [RKLW] .../SQSP AND SQL=18

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4546169 SQL<30

L34 4 .. [WFL] . [VI] [LIMV] [WFLM] . [W] .. [-P] [RKLW] .../SQSP AND SQL<30

=> FIL CAPLUS BIOSIS MEDLINE PCTFULL USPATFULL EUROPATFULL JAPIO SCISEARCH EMBASE

COST IN U.S. DOLLARS SINCE FILE TOTAL ENTRY SESSION

FULL ESTIMATED COST 65.92 998.46

DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS)

SINCE FILE TOTAL
ENTRY SESSION

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L26 L27. (FILE 'HOME' ENTERED AT 14:50:12 ON 04 JAN 2005)

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L2
              2 S WEVLCWTWETCER/SQEP
L3
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L4
L5
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L6
               8 S .{0,99}.WEVLCWTWETCER/SQSP
L7
L8
               6 S WEVLCWTWETCER. {0,99}./SQSP
1.9
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L10
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L11
L12
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L14
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L15
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L16
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L17
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L18
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L19
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L20
              7 S L16
L21
L22
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L23
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L25
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FILE 'CAPLUS, BIOSIS, MEDLINE, PCTFULL, USPATFULL, JAPIO, SCISEARCH,

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2424 S .. [WFL] . [VI] [LIMV] [WFLM] . [W] .. [-P] [RKLW] .../SQSP — Claim

	EMBASE' ENTERED AT 15:31:55 ON 04 JAN 2005				
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L29	0 S L28 AND DUP REM				
L30	224 DUP REM L28 (7 DUPLICATES REMOVED)				
L31	0 S L30 AND (FACTOR (W) VII)				
L32	1 S L30 AND (VII)				
	FILE 'REGISTRY' ENTERED AT 15:44:46 ON 04 JAN 2005				
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L34	4 S [WFL] . [VI] [LIMV] [WFLM] . [W] [-P] [RKLW]/SQSP AND SQL<30				
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	EMBASE! ENTERED AT 15:46:57 ON 04 JAN 2005				

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